

9	18	27	36	45	54
5' GCT CCC CGC GGC TGG CGT CGG GAA AGT ACA GTA AAA AGT CCG AGT GCA GCC					
63	72	81	90	99	108
GCC GGG CGC AGG ATG GGA TCC GGC TCC TCC AGC TAC CGG CCC AAG GCC ATC TAC M G S G S S S Y R P K A I Y					
117	126	135	144	153	162
CTG GAC ATC GAT GGA CGC ATT CAG AAG GTA ATC TTC AGC AAG TAC TGC AAC TCC L D I D G R I Q K V I F S K Y C N S					
171	180	189	198	207	216
AGC GAC ATC ATG GAC CTG TTC TGC ATC GCC ACC GGC CTG CCT CGG AAC ACG ACC S D I M D L F C I A T G L P R N T T					
225	234	243	252	261	270
ATC TCC CTG CTG ACC ACC GAC GAC GCC ATG GTC TCC ATC GAC CCC ACC ATG CCC I S L L T T D D A M V S I D P T M P					
279	288	297	306	315	324
GCG AAT TCA GAA CGC ACT CCG TAC AAA GTG AGA CCT GTG GCC ATC AAG CAA CTC A N S E R T P Y K V R P V A I K Q L					
333	342	351	360	369	378
TCC GCT GGT GTC GAG GAC AAG AGA ACC ACA AGC CGT GGC CAG TCT GCT GAG AGA S A G V E D K R T T S R G Q S A E R					

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FIGURE 1B

TABLE 1

765	774	783	792	801	810
CAC AAG AAG TTG ACT CCT CGA CGC GAT GTT CCC ACT TAC CCC AAG TAC CTG CTC					
H K K L T P R R D V P T Y P K Y L L					
819	828	837	846	855	864
TCT CCA GAG ACC ATC GAG GCC CTG CGG AAG CCG ACC TTT GAC GTC TGG CTT TGG					
S P E T I E A L R K P T F D V W L W					
873	882	891	900	909	918
GAG CCC AAT GAG ATG CTG AGC TGC CTG GAG CAC ATG TAC CAC GAC CTC GGG CTG					
E P N E M L S C L E H M Y H D L G L					
927	936	945	954	963	972
GTC AGG GAC TTC AGC ATC AAC CCT GTC ACC CTC AGG AGG TGG CTG TTC TGT GTC					
V R D F S I N P V T L R R W L F C V					
981	990	999	1008	1017	1026
CAC GAC AAC TAC AGA AAC AAC CCC TTC CAC AAC TTC CGG CAC TGC TTC TGC GTG					
H D N Y R N N P F H N F R H C F C V					
1035	1044	1053	1062	1071	1080
GCC CAG ATG ATG TAC AGC ATG GTC TGG CTC TGC AGT CTC CAG GAG AAG TTC TCA					
A Q M M Y S M V W L C S L Q E K F S					
1089	1098	1107	1116	1125	1134
CAA ACG GAT ATC CTG ATC CTA ATG ACA GCG GCC ATC TGC CAC GAT CTG GAC CAT					
Q T D I L I L M T A A I C H D L D H					

FIGURE 1C

## TABLE 14-600

1143	1152	1161	1170	1179	1188
CCC GGC TAC AAC AAC ACG TAC CAG ATC AAT GCC CGC ACA GAG CTG GCG GTC CGC					
P G Y N N T Y Q I N A R T E L A V R					
1197	1206	1215	1224	1233	1242
TAC AAT GAC ATC TCA CCG CTG GAG AAC CAC CAC TGC GCC GTG GCC TTC CAG ATC					
Y N D I S P L E N H H C A V A F Q I					
1251	1260	1269	1278	1287	1296
CTC GCC GAG CCT GAG TGC AAC ATC TTC TCC AAC ATC CCA CCT GAT GGG TTC AAG					
L A E P E C N I F S N I P P D G F K					
1305	1314	1323	1332	1341	1350
CAG ATC CGA CAG GGA ATG ATC ACA TTA ATC TTG GCC ACT GAC ATG GCA AGA CAT					
Q I R Q G M I T L I L A T D M A R H					
1359	1368	1377	1386	1395	1404
GCA GAA ATT ATG GAT TCT TTC AAA GAG AAA ATG GAG AAT TTT GAC TAC AGC AAC					
A E I M D S F K E K M E N F D Y S N					
1413	1422	1431	1440	1449	1458
GAG GAG CAC ATG ACC CTG CTG AAG ATG ATT TTG ATA AAA TGC TGT GAT ATC TCT					
E E H M T L L K M I L I K C C D I S					
1467	1476	1485	1494	1503	1512
AAC GAG GTC CGT CCA ATG GAA GTC GCA GAG CCT TGG GTG GAC TGT TTA TTA GAG					
N E V R P M E V A E P W V D C L L E					

FIGURE 1D

[illegible]

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TABLE 1. Nucleotide

1899	1908	1917	1926	1935	1944
AGC TGC GCG GGA TCC TTG TGC AGG GAA GAG CTG CCC TGG GCA CCT GGC ACC ACA					
1953	1962	1971	1980	1989	
AGA CCA TGT TTT CTA AGA ACC ATT TTG TTC ACT GAT ACA AAA AAA AAA AA 3'					

FIGURE 1F



118	R	R	E	G	A	F	E	S	G	Q	V	-	-	-	-	-	-	E	P	R	P	R	E	P	Q	G	C	Y	Q	E	G	Q	R	I	P	PDE9A									
200	I	R	H	Y	-	V	S	I	I	R	V	C	N	G	N	K	A	E	K	I	S	E	C	V	Q	S	D	T	R	T	D	N	Q	T	G	K	H	K	D	PDE8A					
103	Q	Q	E	F	D	L	P	S	L	R	V	-	-	-	-	-	-	E	D	N	P	E	L	V	A	A	N	A	A	G	Q	Q	S	A	G	Q	Y	A	g829179						
147	P	E	R	E	E	L	-	I	Q	S	V	L	A	Q	V	A	E	-	-	-	-	-	-	-	Q	F	S	R	A	F	K	I	N	E	L	K	A	E	-	-	V	A	PDE9A		
239	R	R	K	G	S	L	D	V	K	A	V	A	S	R	A	T	E	V	S	S	Q	R	R	H	S	S	M	A	R	I	H	S	M	T	I	E	A	P	I	T	PDE8A				
136	R	S	R	S	P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	R	G	P	P	M	S	Q	I	S	G	V	K	-	-	-	-	-	g829179			
179	N	H	L	A	V	L	E	K	R	V	E	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PDE9A		
279	K	V	I	N	V	I	N	A	A	Q	E	S	S	P	M	P	V	T	E	A	L	D	R	V	L	E	I	L	R	T	T	E	L	Y	S	P	Q	F	G	A	PDE8A				
153	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	g829179			
206	K	K	-	-	-	-	-	-	-	-	-	M	R	E	E	L	A	A	R	S	S	R	T	N	C	P	C	K	Y	S	F	L	D	N	H	K	K	L	T	P	R	R	PDE9A		
319	K	D	D	D	P	H	A	N	D	L	V	G	G	L	M	S	D	G	L	R	R	L	S	G	N	E	Y	V	L	S	T	K	N	T	Q	M	V	S	S	N	PDE8A				
172	-	-	-	-	-	-	-	-	-	-	-	E	T	P	R	E	N	E	L	-	G	T	L	L	G	E	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	g829179	
239	D	V	P	T	Y	P	K	Y	L	L	S	P	E	T	I	E	A	L	R	K	P	T	F	D	V	W	L	W	E	P	N	E	M	-	-	-	-	-	-	-	-	PDE9A			
359	I	I	T	P	I	S	L	D	D	V	P	P	R	I	A	R	A	M	E	N	E	E	Y	-	-	W	D	F	D	I	F	E	L	E	A	A	T	H	N	R	PDE8A				
186	-	-	-	-	-	-	-	-	-	-	-	L	D	T	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	g829179

FIGURE 2B



272	-	L	S	C	L	E	-	H	M	Y	H	D	L	G	L	V	R	D	F	S	I	N	P	V	T	L	R	R	W	L	F	C	V	H	D	N	Y	-	R	N	PDE9A
397	P	L	I	Y	L	G	L	K	M	F	A	R	F	G	I	C	E	F	L	H	C	S	E	S	T	L	R	S	W	L	Q	I	I	E	A	N	Y	H	S	S	PDE8A
204	P	L	T	C	V	A	Y	T	I	F	Q	S	R	E	L	L	T	S	L	M	I	P	P	K	T	F	L	N	F	M	S	T	L	E	D	H	Y	V	K	D	g829179
309	N	P	F	H	N	F	R	H	C	F	C	V	A	Q	M	M	Y	S	M	V	W	L	C	S	L	Q	E	K	F	S	Q	T	D	I	L	I	L	M	T	A	PDE9A
437	N	P	Y	H	N	S	T	H	S	A	D	V	L	H	A	T	A	Y	F	L	S	K	E	R	I	K	E	T	L	D	P	I	D	E	V	A	A	L	I	A	PDE8A
244	N	P	F	H	N	S	L	H	A	A	D	V	T	Q	S	T	N	V	L	L	N	T	P	A	L	E	G	V	F	T	P	L	E	V	G	A	L	F	A	g829179	
349	A	I	C	H	D	L	D	H	P	G	Y	N	N	T	Y	Q	I	N	A	R	T	E	L	A	V	R	Y	N	D	I	S	P	L	E	N	H	H	C	A	V	PDE9A
477	A	T	I	H	D	V	D	H	P	G	R	T	N	S	F	L	C	N	A	G	S	E	L	A	I	L	Y	N	D	T	A	V	L	E	S	H	H	A	A	L	PDE8A
284	A	C	I	H	D	V	D	H	P	G	L	T	N	Q	F	L	V	N	S	S	S	E	L	A	L	M	Y	N	D	E	S	V	L	E	N	H	H	L	A	V	g829179
389	A	F	Q	I	L	A	E	P	E	-	C	N	I	F	S	N	I	P	P	D	G	F	K	Q	I	R	Q	G	M	I	T	L	I	L	A	T	D	M	A	R	PDE9A
517	A	F	Q	L	T	G	D	D	K	-	C	N	I	F	K	N	M	E	R	N	D	Y	R	T	L	R	Q	G	I	I	D	M	V	L	A	T	E	M	T	K	PDE8A
324	A	F	K	L	L	Q	N	Q	G	-	C	D	I	F	C	N	M	Q	K	Q	R	Q	T	L	R	K	M	V	I	D	I	V	L	S	T	D	M	S	K	g829179	
428	H	A	E	I	M	D	S	F	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	E	PDE9A		
557	H	F	E	H	V	N	K	F	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	E	PDE8A		
363	H	M	S	L	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	g829179			

FIGURE 2C

449	H	M	T	L	L	K	M	I	L	I	K	C	C	D	I	S	N	E	V	R	P	M	E	V	A	E	P	W	V	D	C	L	L	E	E	Y	F	M	Q	S	PDE9A
597	N	R	T	L	I	K	R	M	L	I	K	C	A	D	V	S	N	P	C	R	P	L	Q	Y	C	I	E	W	A	A	R	I	S	E	E	Y	F	S	Q	T	PDE8A
392	D	R	I	Q	V	L	E	N	L	V	H	C	A	D	L	S	N	P	T	K	P	L	P	L	Y	K	R	W	V	A	L	L	M	E	E	F	F	L	Q	G	g829179
489	D	R	E	K	S	E	G	L	P	V	A	-	P	F	M	D	R	D	K	V	T	K	A	T	A	Q	I	G	F	I	K	F	V	L	I	P	M	F	E	T	PDE9A
637	D	E	K	Q	Q	G	L	P	V	M	P	-	V	F	D	R	N	T	C	S	I	P	K	S	Q	I	S	F	I	D	Y	F	I	T	D	M	F	D	A	PDE8A	
432	D	K	E	R	E	S	G	M	D	I	S	-	P	M	C	D	R	H	N	A	T	I	E	K	S	Q	V	G	F	I	D	Y	I	V	H	P	L	W	E	T	g829179
528	V	T	K	L	F	P	M	V	E	E	I	M	L	Q	P	L	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PDE9A
677	W	D	A	F	V	-	-	D	L	P	D	L	M	Q	H	L	D	N	N	F	K	Y	W	K	G	L	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PDE8A
471	W	A	S	L	V	H	P	D	A	Q	D	I	L	D	T	L	E	E	N	R	D	Y	Y	Q	S	M	I	P	P	S	P	P	S	G	V	D	E	N	P	g829179	
545	-	E	S	R	D	R	Y	E	-	E	L	K	R	I	D	D	A	-	M	K	E	L	Q	K	K	T	D	S	L	-	-	-	-	-	-	-	-	-	-	-	PDE9A
701	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	D	E	M	K	L	R	N	L	R	P	P	P	E	-	-	-	-	-	-	-	-	-	-	PDE8A
511	Q	E	D	R	I	R	F	Q	V	T	L	E	E	S	D	Q	E	N	L	A	E	L	E	E	G	D	E	S	G	G	E	T	T	T	G	T	T	G	T	g829179	
575	T	E	K	S	R	E	R	S	R	D	-	-	-	-	-	-	-	-	V	K	N	S	E	G	D	C	A	-	-	-	-	-	-	-	-	-	-	-	-	-	PDE9A
713	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	PDE8A	
551	T	A	S	A	L	R	A	G	G	G	G	G	G	G	G	G	M	A	P	R	T	G	C	G	C	Q	N	Q	P	Q	H	G	G	M	-	-	-	-	-	-	g829179

FIGURE 2D

11 g 13

1030630 1120000

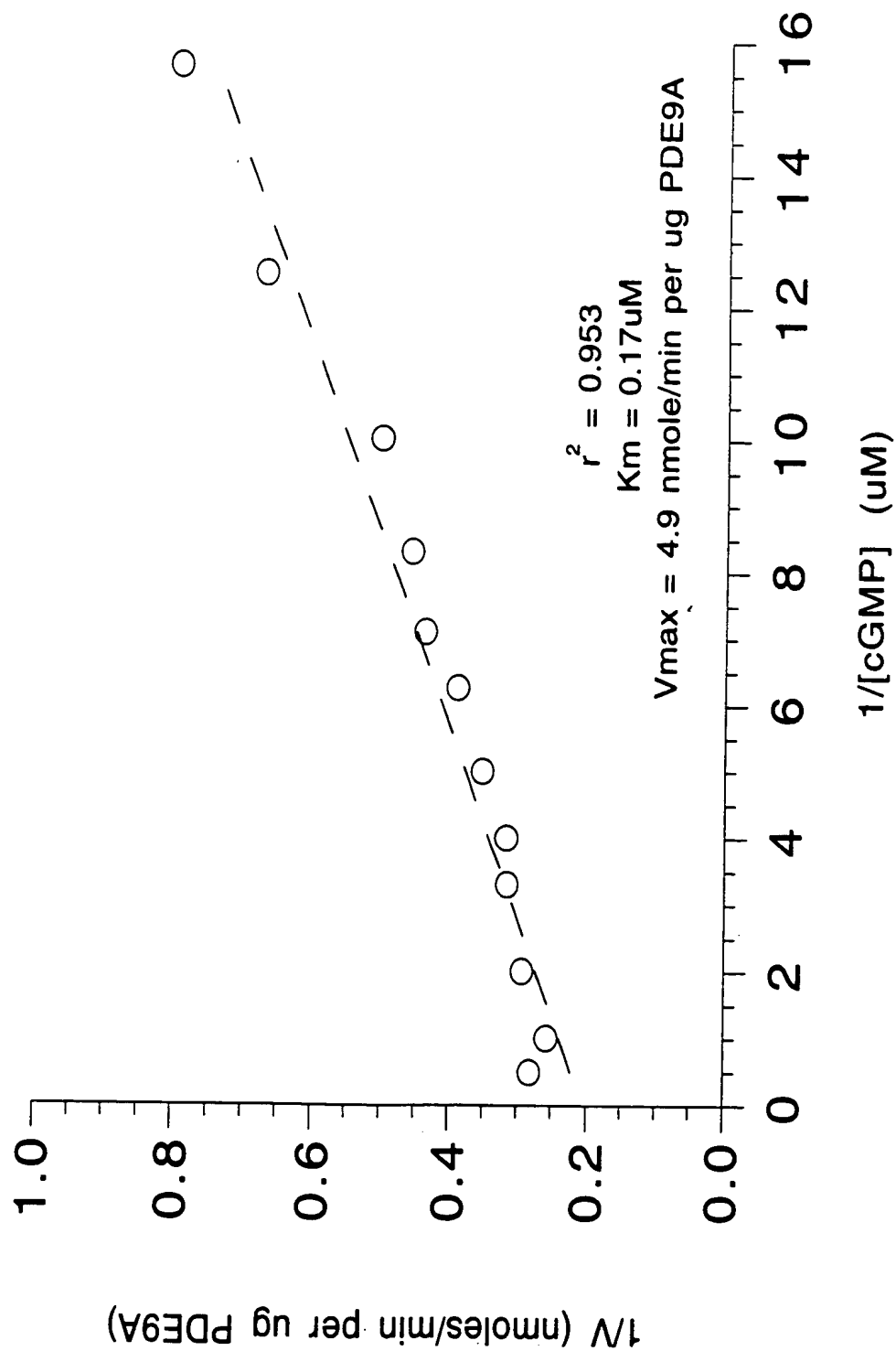


FIGURE 3

TABLE 1

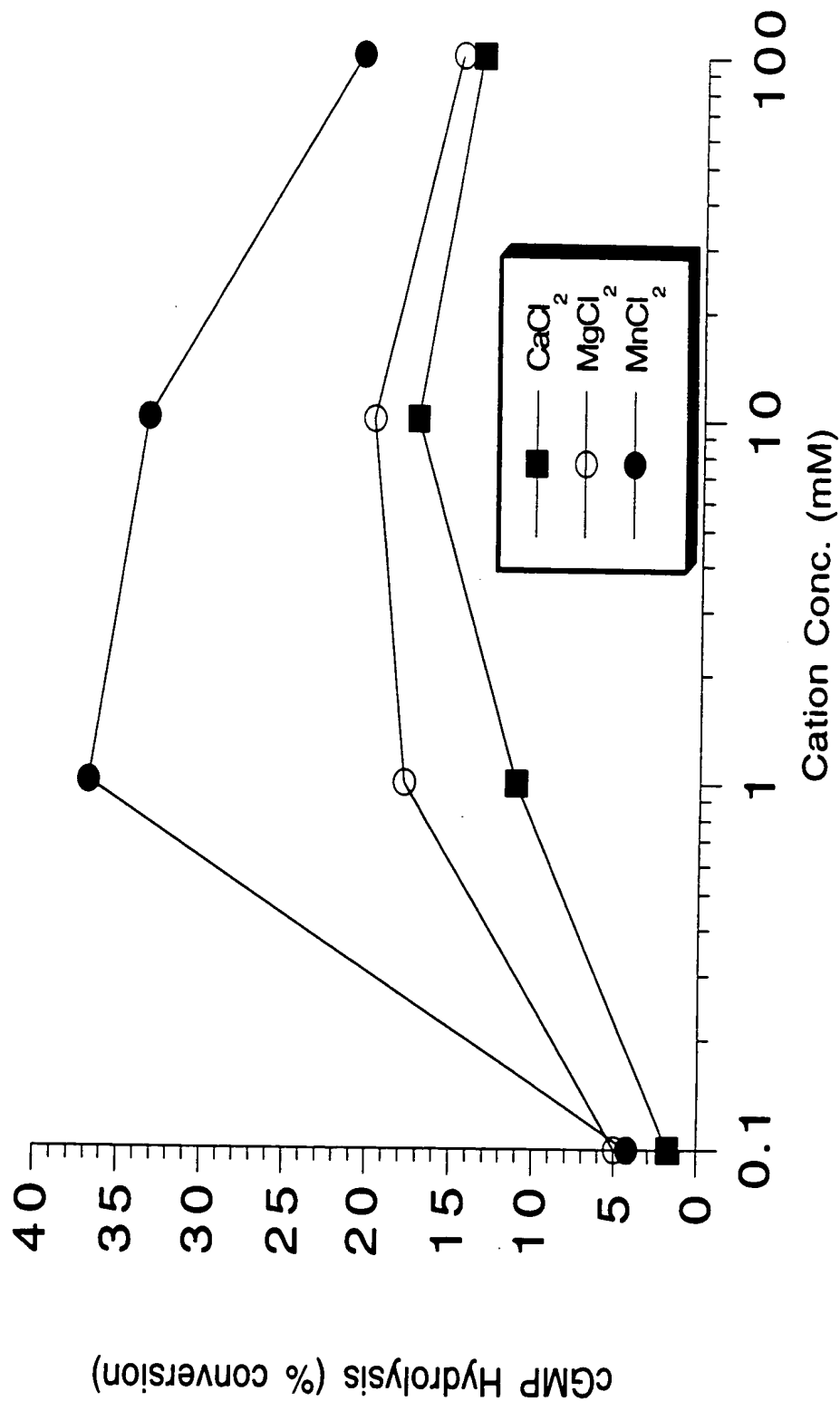


FIGURE 4

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100000 1120000

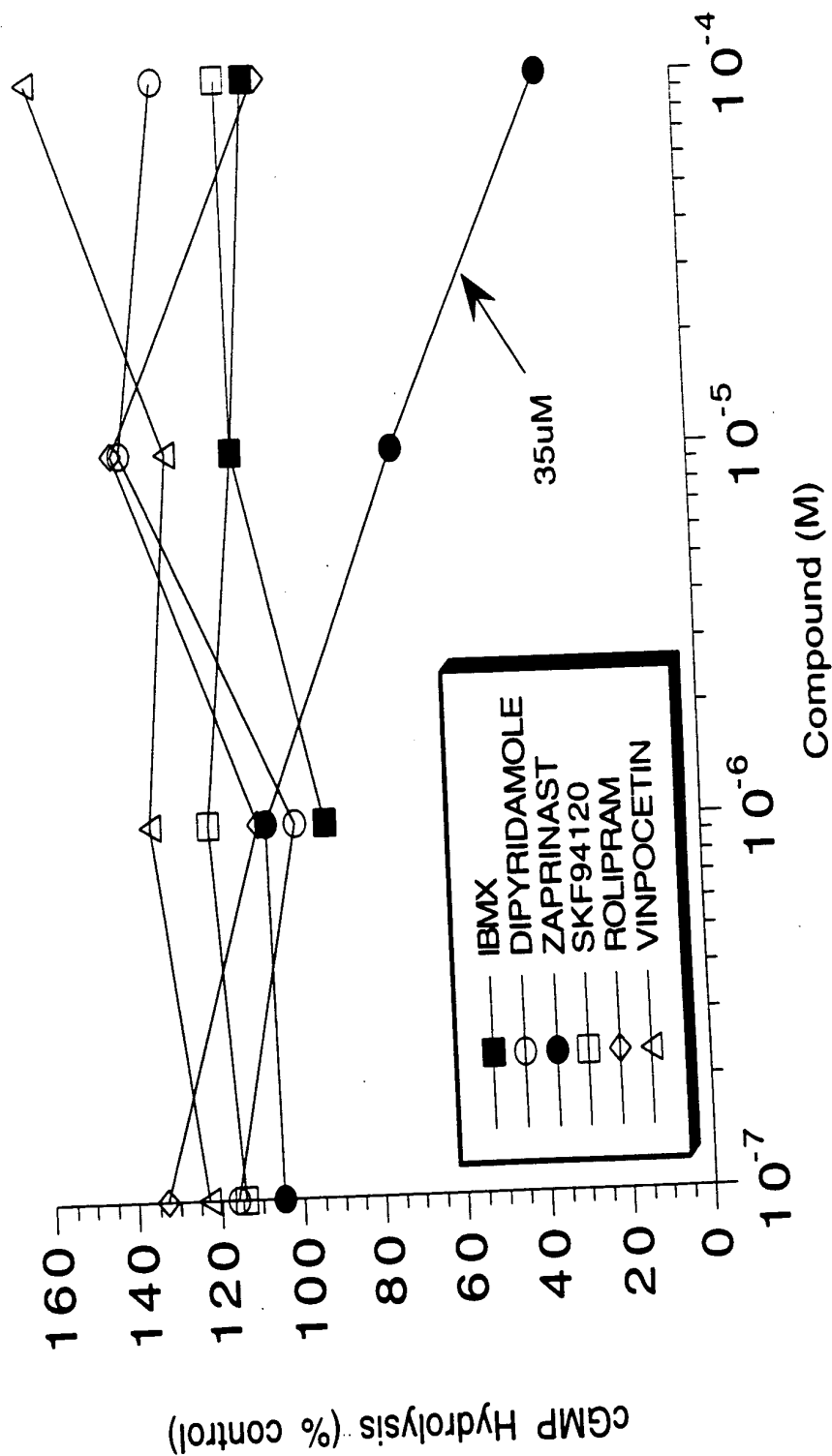


FIGURE 5